

SEQUENCE LISTING

<110> Climent-Johansson, Isabel
Enerback, Sven

<120> PROTEIN COMPLEXES

<130> 13425-102US1

<140> US 10/500,941
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<160> 12

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 501
<212> PRT
<213> Homo sapiens

<400> 1
Met Gln Ala Arg Tyr Ser Val Ser Asp Pro Asn Ala Leu Gly Val Val
1 5 10 15
Pro Tyr Leu Ser Glu Gln Asn Tyr Tyr Arg Ala Ala Gly Ser Tyr Gly
20 25 30
Gly Met Ala Ser Pro Met Gly Val Tyr Ser Gly His Pro Glu Gln Tyr
35 40 45
Ser Ala Gly Met Gly Arg Ser Tyr Ala Pro Tyr His His His Gln Pro
50 55 60
Ala Ala Pro Lys Asp Leu Val Lys Pro Pro Tyr Ser Tyr Ile Ala Leu
65 70 75 80
Ile Thr Met Ala Ile Gln Asn Ala Pro Glu Lys Lys Ile Thr Leu Asn
85 90 95
Gly Ile Tyr Gln Phe Ile Met Asp Arg Phe Pro Phe Tyr Arg Glu Asn
100 105 110
Lys Gln Gly Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Glu
115 120 125
Cys Phe Val Lys Val Pro Arg Asp Asp Lys Lys Pro Gly Lys Gly Ser
130 135 140
Tyr Trp Thr Leu Asp Pro Asp Ser Tyr Asn Met Phe Glu Asn Gly Ser
145 150 155 160
Phe Leu Arg Arg Arg Arg Phe Lys Lys Lys Asp Val Ser Lys Glu
165 170 175
Lys Glu Glu Arg Ala His Leu Lys Glu Pro Pro Pro Ala Ala Ser Lys
180 185 190
Gly Ala Pro Ala Thr Pro His Leu Ala Asp Ala Pro Lys Glu Ala Glu

195	200	205	
Lys Lys Val Val Ile Lys Ser Glu Ala Ala Ser Pro Ala Leu Pro Val			
210	215	220	
Ile Thr Lys Val Glu Thr Leu Ser Pro Glu Ser Ala Leu Gln Gly Ser			
225	230	235	240
Pro Arg Ser Ala Ala Ser Thr Pro Ala Gly Ser Pro Asp Gly Ser Leu			
245	250	255	
Pro Glu His His Ala Ala Ala Pro Asn Gly Leu Pro Gly Phe Ser Val			
260	265	270	
Glu Asn Ile Met Thr Leu Arg Thr Ser Pro Pro Gly Gly Glu Leu Ser			
275	280	285	
Pro Gly Ala Gly Arg Ala Gly Leu Val Val Pro Pro Leu Ala Leu Pro			
290	295	300	
Tyr Ala Ala Ala Pro Pro Ala Ala Tyr Gly Gln Pro Cys Ala Gln Gly			
305	310	315	320
Leu Glu Ala Gly Ala Ala Gly Gly Tyr Gln Cys Ser Met Arg Ala Met			
325	330	335	
Ser Leu Tyr Thr Gly Ala Glu Arg Pro Ala His Met Cys Val Pro Pro			
340	345	350	
Ala Leu Asp Glu Ala Leu Ser Asp His Pro Ser Gly Pro Thr Ser Pro			
355	360	365	
Leu Ser Ala Leu Asn Leu Ala Ala Gly Gln Glu Gly Ala Leu Ala Ala			
370	375	380	
Thr Gly His His His Gln His His Gly His His His Pro Gln Ala Pro			
385	390	395	400
Pro Pro Pro Pro Ala Pro Gln Pro Gln Pro Thr Pro Gln Pro Gly Ala			
405	410	415	
Ala Ala Ala Gln Ala Ala Ser Trp Tyr Leu Asn His Ser Gly Asp Leu			
420	425	430	
Asn His Leu Pro Gly His Thr Phe Ala Ala Gln Gln Gln Thr Phe Pro			
435	440	445	
Asn Val Arg Glu Met Phe Asn Ser His Arg Leu Gly Ile Glu Asn Ser			
450	455	460	
Thr Leu Gly Glu Ser Gln Val Ser Gly Asn Ala Ser Cys Gln Leu Pro			
465	470	475	480
Tyr Arg Ser Thr Pro Pro Leu Tyr Arg His Ala Ala Pro Tyr Ser Tyr			
485	490	495	
Asp Cys Thr Lys Tyr			
500			

<210> 2
 <211> 514
 <212> PRT
 <213> Homo sapiens

<400> 2			
Pro Ser Glu Lys Asn Glu Phe Ser Arg Arg Lys Arg Ser Lys Ser Glu			
1	5	10	15
Asp Met Asp Asn Val Gln Ser Lys Arg Arg Arg Tyr Met Glu Glu Glu			
20	25	30	
Tyr Glu Ala Glu Phe Gln Val Lys Ile Thr Ala Lys Gly Asp Ile Asn			
35	40	45	
Gln Lys Leu Gln Lys Val Ile Gln Trp Leu Leu Glu Glu Lys Leu Cys			
50	55	60	
Ala Leu Gln Cys Ala Val Phe Asp Lys Thr Leu Ala Glu Leu Lys Thr			
65	70	75	80
Arg Val Glu Lys Ile Glu Cys Asn Lys Arg His Lys Thr Val Leu Thr			

85	90	95													
Glu	Leu	Gln	Ala	Lys	Ile	Ala	Arg	Leu	Thr	Lys	Arg	Phe	Glu	Ala	Ala
100								105					110		
Lys	Glu	Asp	Leu	Lys	Lys	Arg	His	Glu	His	Pro	Pro	Asn	Pro	Pro	Val
115								120					125		
Ser	Pro	Gly	Lys	Thr	Val	Asn	Asp	Val	Asn	Ser	Asn	Asn	Asn	Met	Ser
130						135				140					
Tyr	Arg	Asn	Ala	Gly	Thr	Val	Arg	Gln	Met	Leu	Glu	Ser	Lys	Arg	Asn
145						150				155					160
Val	Ser	Glu	Ser	Ala	Pro	Pro	Ser	Phe	Gln	Thr	Pro	Val	Asn	Thr	Val
									165	170				175	
Ser	Ser	Thr	Asn	Leu	Val	Thr	Pro	Pro	Ala	Val	Val	Ser	Ser	Gln	Pro
						180				185				190	
Lys	Leu	Gln	Thr	Pro	Val	Thr	Ser	Gly	Ser	Leu	Thr	Ala	Thr	Ser	Val
						195			200			205			
Leu	Pro	Ala	Pro	Asn	Thr	Ala	Thr	Val	Val	Ala	Thr	Thr	Gln	Val	Pro
						210			215			220			
Ser	Gly	Asn	Pro	Gln	Pro	Thr	Ile	Ser	Leu	Gln	Pro	Leu	Pro	Val	Ile
						225			230			235			240
Leu	His	Val	Pro	Val	Ala	Val	Ser	Ser	Gln	Pro	Gln	Leu	Leu	Gln	Ser
						245			250					255	
His	Pro	Gly	Thr	Leu	Val	Thr	Asn	Gln	Pro	Ser	Gly	Asn	Val	Glu	Phe
						260			265			270			
Ile	Ser	Val	Gln	Ser	Pro	Pro	Thr	Val	Ser	Gly	Leu	Thr	Lys	Asn	Pro
						275			280			285			
Val	Ser	Leu	Pro	Ser	Leu	Pro	Asn	Pro	Thr	Lys	Pro	Asn	Asn	Val	Pro
						290			295			300			
Ser	Val	Pro	Ser	Pro	Ser	Ile	Gln	Arg	Asn	Pro	Thr	Ala	Ser	Ala	Ala
						305			310			315			320
Pro	Leu	Gly	Thr	Thr	Leu	Ala	Val	Gln	Ala	Val	Pro	Thr	Ala	His	Ser
						325			330			335			
Ile	Val	Gln	Ala	Thr	Arg	Thr	Ser	Leu	Pro	Thr	Val	Gly	Pro	Ser	Gly
						340			345			350			
Leu	Tyr	Ser	Pro	Ser	Thr	Asn	Arg	Gly	Pro	Ile	Gln	Met	Lys	Ile	Pro
						355			360			365			
Ile	Ser	Ala	Phe	Ser	Thr	Ser	Ser	Ala	Ala	Glu	Gln	Asn	Ser	Asn	Thr
						370			375			380			
Thr	Pro	Arg	Ile	Glu	Asn	Gln	Thr	Asn	Lys	Thr	Ile	Asp	Ala	Ser	Val
						385			390			395			400
Ser	Lys	Lys	Ala	Ala	Asp	Ser	Thr	Ser	Gln	Cys	Gly	Lys	Ala	Thr	Gly
						405			410			415			
Ser	Asp	Ser	Ser	Gly	Val	Ile	Asp	Leu	Thr	Met	Asp	Asp	Glu	Glu	Ser
						420			425			430			
Gly	Ala	Ser	Gln	Asp	Pro	Lys	Lys	Leu	Asn	His	Thr	Pro	Val	Ser	Thr
						435			440			445			
Met	Ser	Ser	Ser	Gln	Pro	Val	Ser	Arg	Pro	Leu	Gln	Pro	Ile	Gln	Pro
						450			455			460			
Ala	Pro	Pro	Leu	Gln	Pro	Ser	Gly	Val	Pro	Thr	Ser	Gly	Pro	Ser	Gln
						465			470			475			480
Thr	Thr	Ile	His	Leu	Leu	Pro	Thr	Ala	Pro	Thr	Thr	Val	Asn	Val	Thr
						485			490			495			
His	Arg	Pro	Val	Thr	Gln	Val	Thr	Arg	Leu	Pro	Val	Pro	Arg	Ala	
						500			505			510			
Pro	Ala														

<211> 524
 <212> PRT
 <213> Homo sapiens

<400> 3
 Met His Val Glu Thr Gly Pro Asn Gly Glu Gln Ile Arg Lys His Ala
 1 5 10 15
 Gly Gln Lys Arg Thr Tyr Lys Ala Ile Ser Glu Ser Tyr Ala Phe Leu
 20 25 30
 Pro Arg Glu Ala Val Thr Arg Phe Leu Met Ser Cys Ser Glu Cys Gln
 35 40 45
 Lys Arg Met His Leu Asn Pro Asp Gly Thr Asp His Lys Asp Asn Gly
 50 55 60
 Lys Pro Pro Thr Leu Val Thr Ser Met Ile Asp Tyr Asn Met Pro Ile
 65 70 75 80
 Thr Met Ala Tyr Met Lys His Met Lys Leu Gln Leu Leu Asn Ser Gln
 85 90 95
 Gln Asp Glu Asp Glu Ser Ser Ile Glu Ser Asp Glu Phe Asp Met Ser
 100 105 110
 Asp Ser Thr Arg Met Ser Ala Val Asn Ser Asp Leu Ser Ser Asn Leu
 115 120 125
 Glu Glu Arg Met Gln Ser Pro Gln Asn Leu His Gly Gln Gln Asp Asp
 130 135 140
 Asp Ser Ala Ala Glu Ser Phe Asn Gly Asn Glu Thr Leu Gly His Ser
 145 150 155 160
 Ser Ile Ala Ser Gly Gly Thr His Ser Arg Glu Met Gly Asp Ser Asn
 165 170 175
 Ser Asp Gly Lys Thr Gly Leu Glu Gln Asp Glu Gln Pro Leu Asn Leu
 180 185 190
 Ser Asp Ser Pro Leu Ser Ala Gln Leu Thr Ser Glu Tyr Arg Ile Asp
 195 200 205
 Asp His Asn Ser Asn Gly Lys Asn Lys Tyr Lys Asn Leu Leu Ile Ser
 210 215 220
 Asp Leu Lys Met Glu Arg Glu Ala Arg Glu Asn Gly Ser Lys Ser Pro
 225 230 235 240
 Ala His Ser Tyr Ser Ser Tyr Asp Ser Gly Lys Asn Glu Ser Val Asp
 245 250 255
 Arg Gly Ala Glu Asp Leu Ser Leu Asn Arg Gly Asp Glu Asp Glu Asp
 260 265 270
 Asp His Glu Asp His Asp Asp Ser Glu Lys Val Asn Glu Thr Asp Gly
 275 280 285
 Val Glu Ala Glu Arg Leu Lys Ala Phe Asn Met Phe Val Arg Leu Phe
 290 295 300
 Val Asp Glu Asn Leu Asp Arg Met Val Pro Ile Ser Lys Gln Pro Lys
 305 310 315 320
 Glu Lys Ile Gln Ala Ile Ile Asp Ser Cys Arg Arg Gln Phe Pro Glu
 325 330 335
 Tyr Gln Glu Arg Ala Arg Lys Arg Ile Arg Thr Tyr Leu Lys Ser Cys
 340 345 350
 Arg Arg Met Lys Arg Ser Gly Phe Glu Met Ser Arg Pro Ile Pro Ser
 355 360 365
 His Leu Thr Ser Ala Val Ala Glu Ser Ile Leu Ala Ser Ala Cys Glu
 370 375 380
 Ser Glu Ser Arg Asn Ala Ala Lys Arg Met Arg Leu Glu Arg Gln Gln
 385 390 395 400
 Asp Glu Ser Ala Pro Ala Asp Lys Gln Cys Lys Pro Glu Ala Thr Gln
 405 410 415

Ala Thr Tyr Ser Thr Ser Ala Val Pro Gly Ser Gln Asp Val Leu Tyr
 420 425 430
 Ile Asn Gly Asn Gly Thr Tyr Ser Tyr His Ser Tyr Arg Gly Leu Gly
 435 440 445
 Gly Gly Leu Leu Asn Leu Asn Asp Ala Ser Ser Ser Gly Pro Thr Asp
 450 455 460
 Leu Ser Met Lys Arg Gln Leu Ala Thr Ser Ser Gly Ser Ser Ser Ser
 465 470 475 480
 Ser Asn Ser Arg Pro Gln Leu Ser Pro Thr Glu Ile Asn Ala Val Arg
 485 490 495
 Gln Leu Val Ala Gly Tyr Arg Glu Ser Ala Ala Phe Leu Leu Arg Ser
 500 505 510
 Ala Asp Glu Leu Glu Asn Leu Ile Leu Gln Gln Asn
 515 520

<210> 4
 <211> 586
 <212> PRT
 <213> Homo sapiens

<400> 4

Met Asn Pro Thr Asn Thr Val Phe Asp Ala Lys Arg Leu Ile Gly Arg
 1 5 10 15
 Arg Phe Asp Asp Ala Val Val Gln Ser Asp Met Lys His Trp Pro Phe
 20 25 30
 Met Val Val Asn Asp Ala Gly Arg Pro Lys Val Gln Val Glu Tyr Lys
 35 40 45
 Gly Glu Thr Lys Ser Phe Tyr Pro Glu Glu Val Ser Ser Met Val Leu
 50 55 60
 Thr Lys Met Lys Glu Ile Ala Glu Ala Tyr Leu Gly Lys Thr Val Thr
 65 70 75 80
 Asn Ala Val Val Thr Val Pro Ala Tyr Phe Asn Asp Ser Gln Arg Gln
 85 90 95
 Ala Thr Lys Asp Ala Gly Thr Ile Ala Gly Leu Asn Val Leu Arg Ile
 100 105 110
 Ile Asn Glu Pro Thr Ala Ala Ile Ala Tyr Gly Leu Asp Lys Lys
 115 120 125
 Val Gly Ala Glu Arg Asn Val Leu Ile Phe Asp Leu Gly Gly Thr
 130 135 140
 Phe Asp Val Ser Ile Leu Thr Ile Glu Asp Gly Ile Phe Glu Val Lys
 145 150 155 160
 Ser Thr Ala Gly Asp Thr His Leu Gly Gly Glu Asp Phe Asp Asn Arg
 165 170 175
 Met Val Asn His Phe Ile Ala Glu Phe Lys Arg Lys His Lys Lys Asp
 180 185 190
 Ile Ser Glu Asn Lys Arg Ala Val Arg Arg Leu Arg Thr Ala Cys Glu
 195 200 205
 Arg Ala Lys Arg Thr Leu Ser Ser Ser Thr Gln Ala Ser Ile Glu Ile
 210 215 220
 Asp Ser Leu Tyr Glu Gly Ile Asp Phe Tyr Thr Ser Ile Thr Arg Ala
 225 230 235 240
 Arg Phe Glu Glu Leu Asn Ala Asp Leu Phe Arg Gly Thr Leu Asp Pro
 245 250 255
 Val Glu Lys Ala Leu Arg Asp Ala Lys Leu Asp Lys Ser Gln Ile His
 260 265 270
 Asp Ile Val Val Gly Gly Ser Thr Arg Ile Pro Lys Ile Gln Lys
 275 280 285

Leu Leu Gln Asp Phe Phe Asn Gly Lys Glu Leu Asn Lys Ser Ile Asn
 290 295 300
 Pro Asp Glu Ala Val Ala Tyr Gly Ala Ala Val Gln Ala Ala Ile Leu
 305 310 315 320
 Ser Gly Asp Lys Ser Glu Asn Val Gln Asp Leu Leu Leu Asp Val
 325 330 335
 Thr Pro Leu Ser Leu Gly Ile Glu Thr Ala Gly Gly Val Met Thr Val
 340 345 350
 Leu Ile Lys Arg Asn Thr Thr Ile Pro Thr Lys Gln Thr Gln Thr Phe
 355 360 365
 Thr Thr Tyr Ser Asp Asn Gln Pro Gly Val Leu Ile Gln Val Tyr Glu
 370 375 380
 Gly Glu Arg Ala Met Thr Lys Asp Asn Asn Leu Leu Gly Lys Phe Glu
 385 390 395 400
 Leu Thr Gly Ile Pro Pro Ala Pro Arg Gly Val Pro Gln Ile Glu Val
 405 410 415
 Thr Phe Asp Ile Asp Ala Asn Gly Ile Leu Asn Val Ser Ala Val Asp
 420 425 430
 Lys Ser Thr Gly Lys Glu Asn Lys Ile Thr Ile Thr Asn Asp Lys Gly
 435 440 445
 Arg Leu Ser Lys Glu Asp Ile Glu Arg Met Val Gln Glu Ala Glu Lys
 450 455 460
 Tyr Lys Ala Glu Asp Glu Lys Gln Arg Asp Lys Val Ser Ser Lys Asn
 465 470 475 480
 Ser Leu Glu Ser Tyr Ala Phe Asn Met Lys Ala Thr Val Glu Asp Glu
 485 490 495
 Lys Leu Gln Gly Lys Ile Asn Asp Glu Asp Lys Gln Lys Ile Leu Asp
 500 505 510
 Lys Cys Asn Glu Ile Ile Asn Trp Leu Asp Lys Asn Gln Thr Ala Glu
 515 520 525
 Lys Glu Glu Phe Glu His Gln Gln Lys Glu Leu Glu Lys Val Cys Asn
 530 535 540
 Pro Ile Ile Thr Lys Leu Tyr Gln Ser Ala Gly Gly Met Pro Gly Gly
 545 550 555 560
 Met Pro Gly Gly Phe Pro Gly Gly Ala Pro Pro Ser Gly Gly Ala
 565 570 575
 Ser Ser Gly Pro Thr Ile Glu Glu Val Asp
 580 585

<210> 5
 <211> 449
 <212> PRT
 <213> Homo sapiens

<400> 5
 Met Met Leu Ser Thr Glu Gly Arg Glu Gly Phe Val Val Lys Val Arg
 1 5 10 15
 Gly Leu Pro Trp Ser Cys Ser Ala Asp Glu Val Met Arg Phe Phe Ser
 20 25 30
 Asp Cys Lys Ile Gln Asn Gly Thr Ser Gly Ile Arg Phe Ile Tyr Thr
 35 40 45
 Arg Glu Gly Arg Pro Ser Gly Glu Ala Phe Val Glu Leu Glu Ser Glu
 50 55 60
 Glu Glu Val Lys Leu Ala Leu Lys Lys Asp Arg Glu Thr Met Gly His
 65 70 75 80
 Arg Tyr Val Glu Val Phe Lys Ser Asn Ser Val Glu Met Asp Trp Val
 85 90 95

Leu Lys His Thr Gly Pro Asn Ser Pro Asp Thr Ala Asn Asp Gly Phe
 100 105 110
 Val Arg Leu Arg Gly Leu Pro Phe Gly Cys Ser Lys Glu Glu Ile Val
 115 120 125
 Gln Phe Phe Ser Gly Leu Glu Ile Val Pro Asn Gly Met Thr Leu Pro
 130 135 140
 Val Asp Phe Gln Gly Arg Ser Thr Gly Glu Ala Phe Val Gln Phe Ala
 145 150 155 160
 Ser Gln Glu Ile Ala Glu Lys Ala Leu Lys Lys His Lys Glu Arg Ile
 165 170 175
 Gly His Arg Tyr Ile Glu Ile Phe Lys Ser Ser Arg Ala Glu Val Arg
 180 185 190
 Thr His Tyr Asp Pro Pro Arg Lys Leu Met Ala Met Gln Arg Pro Gly
 195 200 205
 Pro Tyr Asp Arg Pro Gly Ala Gly Arg Gly Tyr Asn Ser Ile Gly Arg
 210 215 220
 Gly Ala Gly Phe Glu Arg Met Arg Arg Gly Ala Tyr Gly Gly Gly Tyr
 225 230 235 240
 Gly Gly Tyr Asp Asp Tyr Gly Gly Tyr Asn Asp Gly Tyr Gly Phe Gly
 245 250 255
 Ser Asp Arg Phe Gly Arg Asp Leu Asn Tyr Cys Phe Ser Gly Met Ser
 260 265 270
 Asp His Arg Tyr Gly Asp Gly Ser Ser Phe Gln Ser Thr Thr Gly
 275 280 285
 His Cys Val His Met Arg Gly Leu Pro Tyr Arg Ala Thr Glu Asn Asp
 290 295 300
 Ile Tyr Asn Phe Phe Ser Pro Leu Asn Pro Met Arg Val His Ile Glu
 305 310 315 320
 Ile Gly Pro Asp Gly Arg Val Thr Gly Glu Ala Asp Val Glu Phe Ala
 325 330 335
 Thr His Glu Asp Ala Val Ala Ala Met Ala Lys Asp Lys Ala Asn Met
 340 345 350
 Gln His Arg Tyr Val Glu Leu Phe Leu Asn Ser Thr Ala Gly Thr Ser
 355 360 365
 Gly Gly Ala Tyr Asp His Ser Tyr Val Glu Leu Phe Leu Asn Ser Thr
 370 375 380
 Ala Gly Ala Ser Gly Gly Ala Tyr Gly Ser Gln Met Met Gly Gly Met
 385 390 395 400
 Gly Leu Ser Asn Gln Ser Ser Tyr Gly Gly Pro Ala Ser Gln Gln Leu
 405 410 415
 Ser Gly Gly Tyr Gly Gly Gly Tyr Gly Gly Gln Ser Ser Met Ser Gly
 420 425 430
 Tyr Asp Gln Val Leu Gln Glu Asn Ser Ser Asp Tyr Gln Ser Asn Leu
 435 440 445
 Ala

<210> 6
 <211> 1675
 <212> PRT
 <213> Homo sapiens

<400> 6
 Met Ala Gln Ile Leu Pro Ile Arg Phe Gln Glu His Leu Gln Leu Gln
 1 5 10 15
 Asn Leu Gly Ile Asn Pro Ala Asn Ile Gly Phe Ser Thr Leu Thr Met
 20 25 30

Glu Ser Asp Lys Phe Ile Cys Ile Arg Glu Lys Val Gly Glu Gln Ala
 35 40 45
 Gln Val Val Ile Ile Asp Met Asn Asp Pro Ser Asn Pro Ile Arg Arg
 50 55 60
 Pro Ile Ser Ala Asp Ser Ala Ile Met Asn Pro Ala Ser Lys Val Ile
 65 70 75 80
 Ala Leu Lys Ala Gly Lys Thr Leu Gln Ile Phe Asn Ile Glu Met Lys
 85 90 95
 Ser Lys Met Lys Ala His Thr Met Thr Asp Asp Val Thr Phe Trp Lys
 100 105 110
 Trp Ile Ser Leu Asn Thr Val Ala Leu Val Thr Asp Asn Ala Val Tyr
 115 120 125
 His Trp Ser Met Glu Gly Glu Ser Gln Pro Val Lys Met Phe Asp Arg
 130 135 140
 His Ser Ser Leu Ala Gly Cys Gln Ile Ile Asn Tyr Arg Thr Asp Ala
 145 150 155 160
 Lys Gln Lys Trp Leu Leu Leu Thr Gly Ile Ser Ala Gln Gln Asn Arg
 165 170 175
 Val Val Gly Ala Met Gln Leu Tyr Ser Val Asp Arg Lys Val Ser Gln
 180 185 190
 Pro Ile Glu Gly His Ala Ala Ser Phe Ala Gln Phe Lys Met Glu Gly
 195 200 205
 Asn Ala Glu Glu Ser Thr Leu Phe Cys Phe Ala Val Arg Gly Gln Ala
 210 215 220
 Gly Gly Lys Leu His Ile Ile Glu Val Gly Thr Pro Pro Thr Gly Asn
 225 230 235 240
 Gln Pro Phe Pro Lys Ala Val Asp Val Phe Phe Pro Pro Glu Ala
 245 250 255
 Gln Asn Asp Phe Pro Val Ala Met Gln Ile Ser Glu Lys His Asp Val
 260 265 270
 Val Phe Leu Ile Thr Lys Tyr Gly Tyr Ile His Leu Tyr Asp Leu Glu
 275 280 285
 Thr Gly Thr Cys Ile Tyr Met Asn Arg Ile Ser Gly Glu Thr Ile Phe
 290 295 300
 Val Thr Ala Pro His Glu Ala Thr Ala Gly Ile Ile Gly Val Asn Arg
 305 310 315 320
 Lys Gly Gln Val Leu Ser Val Cys Val Glu Glu Glu Asn Ile Ile Pro
 325 330 335
 Tyr Ile Thr Asn Val Leu Gln Asn Pro Asp Leu Ala Leu Arg Met Ala
 340 345 350
 Val Arg Asn Asn Leu Ala Gly Ala Glu Glu Leu Phe Ala Arg Lys Phe
 355 360 365
 Asn Ala Leu Phe Ala Gln Gly Asn Tyr Ser Glu Ala Ala Lys Val Ala
 370 375 380
 Ala Asn Ala Pro Lys Gly Ile Leu Arg Thr Pro Asp Thr Ile Arg Arg
 385 390 395 400
 Phe Gln Ser Val Pro Ala Gln Pro Gly Gln Thr Ser Pro Leu Leu Gln
 405 410 415
 Tyr Phe Gly Ile Leu Leu Asp Gln Gly Gln Leu Asn Lys Tyr Glu Ser
 420 425 430
 Leu Glu Leu Cys Arg Pro Val Leu Gln Gln Gly Arg Lys Gln Leu Leu
 435 440 445
 Glu Lys Trp Leu Lys Glu Asp Lys Leu Glu Cys Ser Glu Glu Leu Gly
 450 455 460
 Asp Leu Val Lys Ser Val Asp Pro Thr Leu Ala Leu Ser Val Tyr Leu
 465 470 475 480
 Arg Ala Asn Val Pro Asn Lys Val Ile Gln Cys Phe Ala Glu Thr Gly

	485	490	495
Gln Val Gln Lys Ile Val Leu Tyr Ala Lys Lys Val Gly Tyr Thr Pro			
500	505	510	
Asp Trp Ile Phe Leu Leu Arg Asn Val Met Arg Ile Ser Pro Asp Gln			
515	520	525	
Gly Gln Gln Phe Ala Gln Met Leu Val Gln Asp Glu Glu Pro Leu Ala			
530	535	540	
Asp Ile Thr Gln Ile Val Asp Val Phe Met Glu Tyr Asn Leu Ile Gln			
545	550	555	560
Gln Cys Thr Ala Phe Leu Leu Asp Ala Leu Lys Asn Asn Arg Pro Ser			
565	570	575	
Glu Gly Pro Leu Gln Thr Arg Leu Leu Glu Met Asn Leu Met His Ala			
580	585	590	
Pro Gln Val Ala Asp Ala Ile Leu Gly Asn Gln Met Phe Thr His Tyr			
595	600	605	
Asp Arg Ala His Ile Ala Gln Leu Cys Glu Lys Ala Gly Leu Leu Gln			
610	615	620	
Arg Ala Leu Glu His Phe Thr Asp Leu Tyr Asp Ile Lys Arg Ala Val			
625	630	635	640
Val His Thr His Leu Leu Asn Pro Glu Trp Leu Val Asn Tyr Phe Gly			
645	650	655	
Ser Leu Ser Val Glu Asp Ser Leu Glu Cys Leu Arg Ala Met Leu Ser			
660	665	670	
Ala Asn Ile Arg Gln Asn Leu Gln Ile Cys Val Gln Val Ala Ser Lys			
675	680	685	
Tyr His Glu Gln Leu Ser Thr Gln Ser Leu Ile Glu Leu Phe Glu Ser			
690	695	700	
Phe Lys Ser Phe Glu Gly Leu Phe Tyr Phe Leu Gly Ser Ile Val Asn			
705	710	715	720
Phe Ser Gln Asp Pro Asp Val His Phe Lys Tyr Ile Gln Ala Ala Cys			
725	730	735	
Lys Thr Gly Gln Ile Lys Glu Val Glu Arg Ile Cys Arg Glu Ser Asn			
740	745	750	
Cys Tyr Asp Pro Glu Arg Val Lys Asn Phe Leu Lys Glu Ala Lys Leu			
755	760	765	
Thr Asp Gln Leu Pro Leu Ile Ile Val Cys Asp Arg Phe Asp Phe Val			
770	775	780	
His Asp Leu Val Leu Tyr Leu Tyr Arg Asn Asn Leu Gln Lys Tyr Ile			
785	790	795	800
Glu Ile Tyr Val Gln Lys Val Asn Pro Ser Arg Leu Pro Val Val Ile			
805	810	815	
Gly Gly Leu Leu Asp Val Asp Cys Ser Glu Asp Val Ile Lys Asn Leu			
820	825	830	
Ile Leu Val Val Arg Gly Gln Phe Ser Thr Asp Glu Leu Val Ala Glu			
835	840	845	
Val Glu Lys Arg Asn Arg Leu Lys Leu Leu Leu Pro Trp Leu Glu Ala			
850	855	860	
Arg Ile His Glu Gly Cys Glu Glu Pro Ala Thr His Asn Ala Leu Ala			
865	870	875	880
Lys Ile Tyr Ile Asp Ser Asn Asn Pro Glu Arg Phe Leu Arg Glu			
885	890	895	
Asn Pro Tyr Tyr Asp Ser Arg Val Val Gly Lys Tyr Cys Glu Lys Arg			
900	905	910	
Asp Pro His Leu Ala Cys Val Ala Tyr Glu Arg Gly Gln Cys Asp Leu			
915	920	925	
Glu Leu Ile Asn Val Cys Asn Glu Asn Ser Leu Phe Lys Ser Leu Ser			
930	935	940	

Arg Tyr Leu Val Arg Arg Lys Asp Pro Glu Leu Trp Gly Ser Val Leu
 945 950 955 960
 Leu Glu Ser Asn Pro Tyr Arg Arg Pro Leu Ile Asp Gln Val Val Gln
 965 970 975
 Thr Ala Leu Ser Glu Thr Gln Asp Pro Glu Glu Val Ser Val Thr Val
 980 985 990
 Lys Ala Phe Met Thr Ala Asp Leu Pro Asn Glu Leu Ile Glu Leu Leu
 995 1000 1005
 Glu Lys Ile Val Leu Asp Asn Ser Val Phe Ser Glu His Arg Asn Leu
 1010 1015 1020
 Gln Asn Leu Leu Ile Leu Thr Ala Ile Lys Ala Asp Arg Thr Arg Val
 1025 1030 1035 1040
 Met Glu Tyr Ile Asn Arg Leu Asp Asn Tyr Asp Ala Pro Asp Ile Ala
 1045 1050 1055
 Asn Ile Ala Ile Ser Asn Glu Leu Phe Glu Glu Ala Phe Ala Ile Phe
 1060 1065 1070
 Arg Lys Phe Asp Val Asn Thr Ser Ala Val Gln Val Leu Ile Glu His
 1075 1080 1085
 Ile Gly Asn Leu Asp Arg Ala Tyr Glu Phe Ala Glu Arg Cys Asn Glu
 1090 1095 1100
 Pro Ala Val Trp Ser Gln Leu Ala Lys Ala Gln Leu Gln Lys Gly Met
 1105 1110 1115 1120
 Val Lys Glu Ala Ile Asp Ser Tyr Ile Lys Ala Asp Asp Pro Ser Ser
 1125 1130 1135
 Tyr Met Glu Val Val Gln Ala Ala Asn Thr Ser Gly Asn Trp Glu Glu
 1140 1145 1150
 Leu Val Lys Tyr Leu Gln Met Ala Arg Lys Lys Ala Arg Glu Ser Tyr
 1155 1160 1165
 Val Glu Thr Glu Leu Ile Phe Ala Leu Ala Lys Thr Asn Arg Leu Ala
 1170 1175 1180
 Glu Leu Glu Glu Phe Ile Asn Gly Pro Asn Asn Ala His Ile Gln Gln
 1185 1190 1195 1200
 Val Gly Asp Arg Cys Tyr Asp Glu Lys Met Tyr Asp Ala Ala Lys Leu
 1205 1210 1215
 Leu Tyr Asn Asn Val Ser Asn Phe Gly Arg Leu Ala Ser Thr Leu Val
 1220 1225 1230
 His Leu Gly Glu Tyr Gln Ala Ala Val Asp Gly Ala Arg Lys Ala Asn
 1235 1240 1245
 Ser Thr Arg Thr Trp Lys Glu Val Cys Phe Ala Cys Val Asp Gly Lys
 1250 1255 1260
 Glu Phe Arg Leu Ala Gln Met Cys Gly Leu His Ile Val Val His Ala
 1265 1270 1275 1280
 Asp Glu Leu Glu Leu Ile Asn Tyr Tyr Gln Asp Arg Gly Tyr Phe
 1285 1290 1295
 Glu Glu Leu Ile Thr Met Leu Glu Ala Ala Leu Gly Leu Glu Arg Ala
 1300 1305 1310
 His Met Gly Met Phe Thr Glu Leu Ala Ile Leu Tyr Ser Lys Phe Lys
 1315 1320 1325
 Pro Gln Lys Met Arg Glu His Leu Glu Leu Phe Trp Ser Arg Val Asn
 1330 1335 1340
 Ile Pro Lys Val Leu Arg Ala Ala Glu Gln Ala His Leu Trp Ala Glu
 1345 1350 1355 1360
 Leu Val Phe Leu Tyr Asp Lys Tyr Glu Glu Tyr Asp Asn Ala Ile Ile
 1365 1370 1375
 Thr Met Met Asn His Pro Thr Asp Ala Trp Lys Glu Gly Gln Phe Lys
 1380 1385 1390
 Asp Ile Ile Thr Lys Val Ala Asn Val Glu Leu Tyr Tyr Arg Ala Ile

1395	1400	1405	
Gln Phe Tyr Leu Glu Phe Lys Pro Leu Leu Leu Asn Asp Leu Leu Met			
1410	1415	1420	
Val Leu Ser Pro Arg Leu Asp His Thr Arg Ala Val Asn Tyr Phe Ser			
1425	1430	1435	1440
Lys Val Lys Gln Leu Pro Leu Val Lys Pro Tyr Leu Arg Ser Val Gln			
1445	1450	1455	
Asn His Asn Asn Lys Ser Val Asn Glu Ser Leu Asn Asn Leu Phe Ile			
1460	1465	1470	
Thr Glu Glu Asp Tyr Gln Ala Leu Arg Thr Ser Ile Asp Ala Tyr Asp			
1475	1480	1485	
Asn Phe Asp Asn Ile Ser Leu Ala Gln Arg Leu Glu Lys His Glu Leu			
1490	1495	1500	
Ile Glu Phe Arg Arg Ile Ala Ala Tyr Leu Phe Lys Gly Asn Asn Arg			
1505	1510	1515	1520
Trp Lys Gln Ser Val Glu Leu Cys Lys Lys Asp Ser Leu Tyr Lys Asp			
1525	1530	1535	
Ala Met Gln Tyr Ala Ser Glu Ser Lys Asp Thr Glu Leu Ala Glu Glu			
1540	1545	1550	
Leu Leu Gln Trp Phe Leu Gln Glu Glu Lys Arg Glu Cys Phe Gly Ala			
1555	1560	1565	
Cys Leu Phe Thr Cys Tyr Asp Leu Leu Arg Pro Asp Val Val Leu Glu			
1570	1575	1580	
Thr Ala Trp Arg His Asn Ile Met Asp Phe Ala Met Pro Tyr Phe Ile			
1585	1590	1595	1600
Gln Val Met Lys Glu Tyr Leu Thr Lys Val Asp Lys Leu Asp Ala Ser			
1605	1610	1615	
Glu Ser Leu Arg Lys Glu Glu Glu Gln Ala Thr Glu Thr Gln Pro Ile			
1620	1625	1630	
Val Tyr Gly Gln Pro Gln Leu Met Leu Thr Ala Gly Pro Ser Val Ala			
1635	1640	1645	
Val Pro Pro Gln Ala Pro Phe Gly Tyr Gly Tyr Thr Ala Pro Pro Tyr			
1650	1655	1660	
Gly Gln Pro Gln Pro Gly Phe Gly Tyr Ser Met			
1665	1670	1675	

<210> 7
 <211> 903
 <212> PRT
 <213> Homo sapiens

<400> 7			
Met Ala Ile Gln Phe Arg Ser Leu Phe Pro Leu Ala Leu Pro Gly Met			
1	5	10	15
Leu Ala Leu Leu Gly Trp Trp Trp Phe Phe Ser Arg Lys Lys Gly His			
20	25	30	
Val Ser Ser His Asp Glu Gln Gln Val Glu Ala Gly Ala Val Gln Leu			
35	40	45	
Arg Ala Asp Pro Ala Ile Lys Glu Pro Leu Pro Val Glu Asp Val Cys			
50	55	60	
Pro Lys Val Val Ser Thr Pro Pro Ser Val Thr Glu Pro Pro Glu Lys			
65	70	75	80
Glu Leu Ser Thr Val Ser Lys Leu Pro Ala Glu Pro Pro Ala Leu Leu			
85	90	95	
Gln Thr His Pro Pro Cys Arg Arg Ser Glu Ser Ser Gly Ile Leu Pro			
100	105	110	
Asn Thr Thr Asp Met Arg Leu Arg Pro Gly Thr Arg Arg Asp Asp Ser			

115	120	125
Thr Lys Leu Glu Leu Ala Leu Thr Gly Gly Glu Ala Lys Ser Ile Pro		
130	135	140
Leu Glu Cys Pro Leu Ser Ser Pro Lys Gly Val Leu Phe Ser Ser Lys		
145	150	155
Ser Ala Glu Val Cys Lys Gln Asp Ser Pro Phe Ser Arg Val Pro Arg		
165	170	175
Lys Val Gln Pro Gly Tyr Pro Val Val Pro Ala Glu Lys Arg Ser Ser		
180	185	190
Gly Glu Arg Ala Arg Glu Thr Gly Gly Ala Glu Gly Thr Gly Asp Ala		
195	200	205
Val Leu Gly Glu Lys Val Leu Glu Glu Ala Leu Leu Ser Arg Glu His		
210	215	220
Val Leu Glu Leu Glu Asn Ser Lys Gly Pro Ser Leu Ala Ser Leu Glu		
225	230	235
Gly Glu Glu Asp Lys Gly Lys Ser Ser Ser Gln Val Val Gly Pro		
245	250	255
Val Gln Glu Glu Glu Tyr Val Ala Glu Lys Leu Pro Ser Arg Phe Ile		
260	265	270
Glu Ser Ala His Thr Glu Leu Ala Lys Asp Asp Ala Ala Pro Ala Pro		
275	280	285
Pro Val Ala Asp Ala Lys Ala Gln Asp Arg Gly Val Glu Gly Glu Leu		
290	295	300
Gly Asn Glu Glu Ser Leu Asp Arg Asn Glu Glu Gly Leu Asp Arg Asn		
305	310	315
Glu Glu Gly Leu Asp Arg Asn Glu Glu Ser Leu Asp Arg Asn Glu Glu		
325	330	335
Gly Leu Asp Arg Asn Glu Glu Ile Lys Arg Ala Ala Phe Gln Ile Ile		
340	345	350
Ser Gln Val Ile Ser Glu Ala Thr Glu Gln Val Leu Ala Thr Thr Val		
355	360	365
Gly Lys Val Ala Gly Arg Val Cys Gln Ala Ser Gln Leu Gln Gly Gln		
370	375	380
Lys Glu Glu Ser Cys Val Pro Val His Gln Lys Thr Val Leu Gly Pro		
385	390	395
Asp Thr Ala Glu Pro Ala Thr Ala Glu Ala Ala Val Ala Pro Pro Asp		
405	410	415
Ala Gly Leu Pro Leu Pro Gly Leu Pro Ala Glu Gly Ser Pro Pro Pro		
420	425	430
Lys Thr Tyr Val Ser Cys Leu Lys Ser Leu Leu Ser Ser Pro Thr Lys		
435	440	445
Asp Ser Lys Pro Asn Ile Ser Ala His His Ile Ser Leu Ala Ser Cys		
450	455	460
Leu Ala Leu Thr Thr Pro Ser Glu Glu Leu Pro Asp Arg Ala Gly Ile		
465	470	475
Leu Val Glu Asp Ala Thr Cys Val Thr Cys Met Ser Asp Ser Ser Gln		
485	490	495
Ser Val Pro Leu Val Ala Ser Pro Gly His Cys Ser Asp Ser Phe Ser		
500	505	510
Thr Ser Gly Leu Glu Asp Ser Cys Thr Glu Thr Ser Ser Ser Pro Arg		
515	520	525
Asp Lys Ala Ile Thr Pro Pro Leu Pro Glu Ser Thr Val Pro Phe Ser		
530	535	540
Asn Gly Val Leu Lys Gly Glu Leu Ser Asp Leu Gly Ala Glu Asp Gly		
545	550	555
Trp Thr Met Asp Ala Glu Ala Asp His Ser Gly Gly Ser Asp Arg Asn		
565	570	575

Ser Met Asp Ser Val Asp Ser Cys Cys Ser Leu Lys Lys Thr Glu Ser
 580 585 590
 Phe Gln Asn Ala Gln Ala Gly Ser Asn Pro Lys Lys Val Asp Leu Ile
 595 600 605
 Ile Trp Glu Ile Glu Val Pro Lys His Leu Val Gly Arg Leu Ile Gly
 610 615 620
 Lys Gln Gly Arg Tyr Val Ser Phe Leu Lys Gln Thr Ser Gly Ala Lys
 625 630 635 640
 Ile Tyr Ile Ser Thr Leu Pro Tyr Thr Gln Ser Val Gln Ile Cys His
 645 650 655
 Ile Glu Gly Ser Gln His His Val Asp Lys Ala Leu Asn Leu Ile Gly
 660 665 670
 Lys Lys Phe Lys Glu Leu Asn Leu Thr Asn Ile Tyr Ala Pro Pro Leu
 675 680 685
 Pro Ser Leu Ala Leu Pro Ser Leu Pro Met Thr Ser Trp Leu Met Leu
 690 695 700
 Pro Asp Gly Ile Thr Val Glu Val Ile Val Val Asn Gln Val Asn Ala
 705 710 715 720
 Gly His Leu Phe Val Gln Gln His Thr His Pro Thr Phe His Ala Leu
 725 730 735
 Arg Ser Leu Asp Gln Gln Met Tyr Leu Cys Tyr Ser Gln Pro Gly Ile
 740 745 750
 Pro Thr Leu Pro Thr Pro Val Glu Ile Thr Val Ile Cys Ala Ala Pro
 755 760 765
 Gly Ala Asp Gly Ala Trp Trp Arg Ala Gln Val Val Ala Ser Tyr Glu
 770 775 780
 Glu Thr Asn Glu Val Glu Ile Arg Tyr Val Asp Tyr Gly Gly Tyr Lys
 785 790 795 800
 Arg Val Lys Val Asp Val Leu Arg Gln Ile Arg Ser Asp Phe Val Thr
 805 810 815
 Leu Pro Phe Gln Gly Ala Glu Val Leu Leu Asp Ser Val Met Pro Leu
 820 825 830
 Ser Asp Asp Asp Gln Phe Ser Pro Glu Ala Asp Ala Ala Met Ser Glu
 835 840 845
 Met Thr Gly Asn Thr Ala Leu Leu Ala Gln Val Thr Ser Tyr Ser Pro
 850 855 860
 Thr Gly Leu Pro Leu Ile Gln Leu Trp Ser Val Val Gly Asp Glu Val
 865 870 875 880
 Val Leu Ile Asn Arg Ser Leu Val Glu Arg Gly Leu Ala Gln Trp Val
 885 890 895
 Asp Ser Tyr Tyr Thr Ser Leu
 900

<210> 8
 <211> 197
 <212> PRT
 <213> Homo sapiens

<400> 8
 Met Met Phe Pro Gln Ser Arg His Ser Gly Ser Ser His Leu Pro Gln
 1 5 10 15
 Gln Leu Lys Phe Thr Thr Ser Asp Ser Cys Asp Arg Ile Thr Asp Glu
 20 25 30
 Phe Gln Leu Leu Gln Ala Gln Tyr His Ser Leu Lys Leu Glu Cys Asp
 35 40 45
 Lys Leu Ala Ser Glu Lys Ser Glu Met Gln Arg His Tyr Val Met Tyr
 50 55 60

Tyr Glu Met Ser Tyr Gly Leu Asn Ile Glu Met His Lys Gln Ala Glu
 65 70 75 80
 Ile Val Lys Arg Leu Asn Gly Ile Cys Ala Gln Val Leu Pro Tyr Leu
 85 90 95
 Ser Gln Glu His Gln Gln Gln Val Leu Gly Ala Ile Glu Arg Ala Lys
 100 105 110
 Gln Val Thr Ala Pro Glu Leu Asn Ser Ile Ile Arg Gln Gln Leu Gln
 115 120 125
 Ala His Gln Leu Ser Gln Leu Gln Ala Leu Ala Leu Pro Leu Thr Pro
 130 135 140
 Leu Pro Val Gly Leu Gln Pro Pro Ser Leu Pro Ala Val Ser Ala Gly
 145 150 155 160
 Thr Gly Leu Leu Ser Leu Ser Ala Leu Gly Ser Gln Ala His Leu Ser
 165 170 175
 Lys Glu Asp Lys Asn Gly His Asp Gly Asp Thr His Gln Glu Asp Asp
 180 185 190
 Gly Glu Lys Ser Asp
 195

<210> 9
 <211> 33
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> PCR primer

<400> 9
 cgccggatccc catgcaggcg cgctactccg tgt 33

<210> 10
 <211> 33
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> PCR primer

<400> 10
 cgcggatcct cagtatttcg tgcagtcgta gga 33

<210> 11
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> PCR primer

<400> 11
 taatacgact cactataggg 20

<210> 12
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
<223> PCR primer

<400> 12
aggcggtgaa tgtaagcgt